L14

	(FILE 'HOME' ENTERED AT 14:25:29 ON 07 JAN 2008)
Ll	FILE 'REGISTRY' ENTERED AT 14:25:37 ON 07 JAN 2008 296092 S RKR/SQSP
L2 L3 L4	· · · · · · · · · · · · · · · · · · ·
L5	FILE 'REGISTRY' ENTERED AT 14:28:47 ON 07 JAN 2008 0 S RKR/SPSP AND PEG
	FILE 'CAPLUS, MEDLINE' ENTERED AT 15:12:37 ON 07 JAN 2008
L6	338 S BRANCHED (P) (LYS OR LYSINE) (P) (ARG OR ARGININE)
L7	4 S L6 (P) RESIN
L8	2 S L6 (P) IMMOBILIZ?
L9	12 S L6 (P) AFFINITY
L10	18 S L7 OR L8 OR L9
L11	
L12	1 S L11 AND (RESIN OR AFFINITY OR IMMOBILIZ? OR DENDRIMER)
	FILE 'CAPLUS, MEDLINE' ENTERED AT 15:20:13 ON 07 JAN 2008
L13	
T 1 /	ρε ε τιρ λωρ (ρεετώ ορ λεετώτην ορ τωμορτίτος ορ ρερήτος)

25 S L13 AND (RESIN OR AFFINITY OR IMMOBILIZ? OR PEPTIDE)

## (FILE 'HOME' ENTERED AT 09:54:11 ON 07 JAN 2008)

FILE 'REGISTRY' ENTERED AT 09:54:22 ON 07 JAN 2008 L1 5 S RKR/SQEP

FILE 'CAPLUS, BIOSIS' ENTERED AT 09:54:41 ON 07 JAN 2008 L2 3 S L1

L2	3 8	3 L1
	FILE 'CAPLUS	, BIOSIS' ENTERED AT 09:59:47 ON 07 JAN 2008
L3	166982	POLYSTYRENE
L4	165846	G (POLYETHYLENE (A) GLYCOL) OR PEG
L5	20171 \$	G TRIPEPTIDE
L6	2 3	5 L3 (P) L4 (P) L5
L7	4 5	G L3 AND L4 AND L5
L8		S L7 NOT L6
L9	259332	G (ARGININE OR ARG)
L10	34 9	L9 AND L3 AND L4
L11	24 5	L10 AND (RESIN OR IMMOBILIZ? OR AFFINITY)
L12	20 \$	S L11 AND PEPTIDE
L13	1632 8	S L9 AND (PEPTIDE (P) COLUMN)
L14		S L13 AND AFFINITY
L15	343 8	S L9 AND (PEPTIDE (P) COLUMN (P) AFFINITY)
L16	121 8	S L15 AND (LYS OR LYSINE)
L17	0 5	G L16 AND L3 AND L4
L18		S L16 AND RESIN
L19	415 8	RAPP W?/AU
L20	6 8	S L19 AND AFFINITY AND POLYSTYRENE
L21	173 8	G DEPPISCH R?/AU
L22		L21 AND AFFINITY AND POLYSTYRENE NOT L20
L23	0 5	GOHL H?/AU AND AFFINITY AND POLYSTYRENE NOT L20
L24		S WITTNER B?/AU
L25		WITTNER B?/AU AND AFFINITYA AND POLYSTYRENE NOT L20
L26		WITTNER B?/AU AND AFFINITY AND POLYSTYRENE NOT L20
L27	1 8	BECK W?/AU AND AFFINITY AND POLYSTYRENE NOT L20

FILE 'REGISTRY' ENTERED AT 16:05:58 ON 07 JAN 2008 L15 STRUCTURE UPLOADED

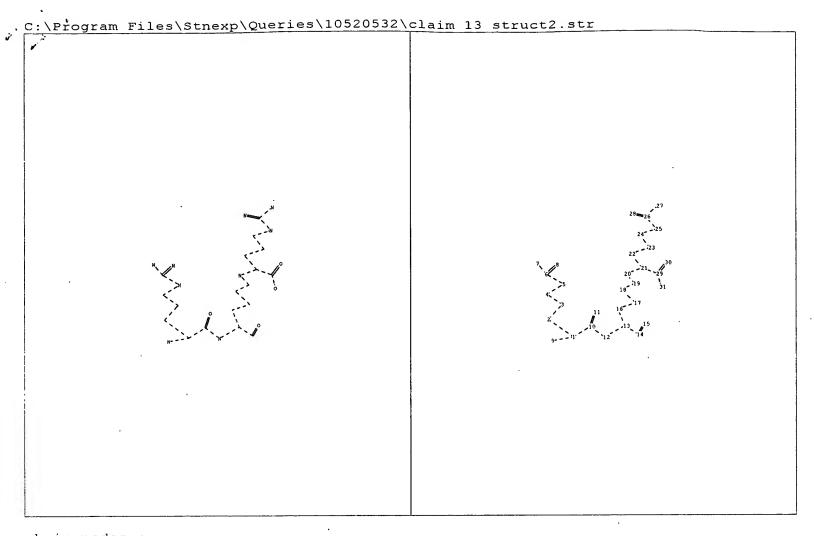
FILE 'CAPLUS' ENTERED AT 16:06:37 ON 07 JAN 2008

FILE 'REGISTRY' ENTERED AT 16:06:51 ON 07 JAN 2008

0 S L15 SSS SAM

L17 0 S L15 SSS FULL

L16



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chain nodes :
    1  2  3  4  5  6  7  8  9  10  11  12  13  14  15  16  17  18  19  20  21
    22  23  24  25  26  27  28  29  30  31

chain bonds :
    1-2  1-9  1-10  2-3  3-4  4-5  5-6  6-7  6-8  10-11  10-12  12-13  13-14
    13-16  14-15  16-17  17-18  18-19  19-20  20-21  21-22  21-29  22-23  23-24
    24-25  25-26  26-27  26-28  29-30  29-31

exact/norm bonds :
    1-2  1-9  1-10  2-3  3-4  4-5  5-6  6-7  6-8  10-11  10-12  12-13  13-14
    13-16  14-15  16-17  17-18  18-19  19-20  20-21  21-22  21-29  22-23  23-24
    24-25  25-26  26-27  26-28  29-30  29-31
```

```
Match level:
1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS 20:CLASS 21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS 31:CLASS
```